

F I G. 1

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CANDIDATE SEQUENCE1: 0000

LIKELIHOOD OF

CANDIDATE SEQUENCE1: 31

CANDIDATE SEQUENCE2: 0101

LIKELIHOOD OF

CANDIDATE SEQUENCE2: 1

CANDIDATE SEQUENCE3: 0110

LIKELIHOOD OF

CANDIDATE SEQUENCE3: 5

CANDIDATE SEQUENCE4: 0011

LIKELIHOOD OF

CANDIDATE SEQUENCE4: 3

RELATIVE : 31-(1+5+3)/3

= 28

28>15 : THRESHOLD VALUE

**MODULATION** 

SCHEME : HIGH DATA RATE

CODING RATE: HIGH

TRANSMISSION

POWER : OMIT

CANDIDATE SEQUENCE1: 0000

LIKELIHOOD OF

CANDIDATE SEQUENCE1: 15

CANDIDATE SEQUENCE2: 0101

LIKELIHOOD OF

CANDIDATE SEQUENCE2: 12

CANDIDATE SEQUENCE3: 0110

LIKELIHOOD OF

CANDIDATE SEQUENCE3: 6

CANDIDATE SEQUENCE4: 0011

LIKELIHOOD OF

CANDIDATE SEQUENCE4: 9

RELATIVE : 15-(12+6+9)/3

= 6

6<15 : THRESHOLD VALUE

**MODULATION** 

SCHEME

: LOW DATA RATE

CODING RATE : LOW

**TRANSMISSION** 

POWER

: LARGE

FIG. 2A

F I G. 2B

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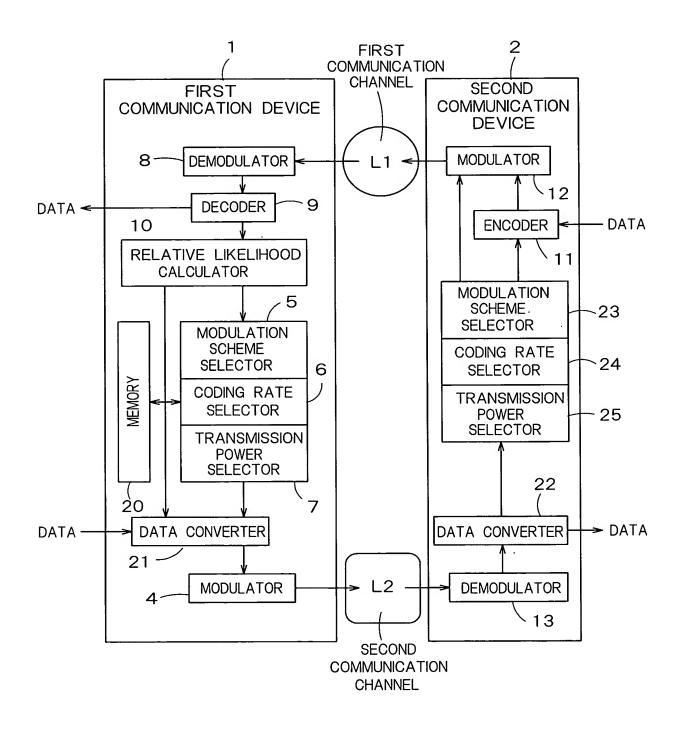


FIG. 3